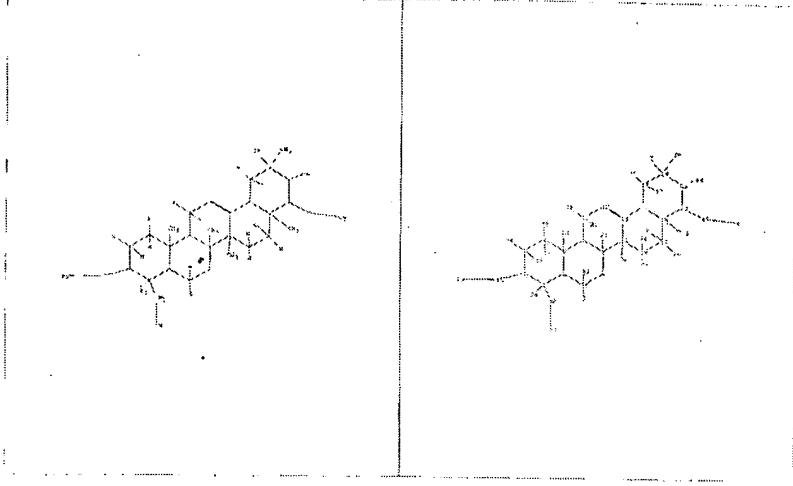


BEST AVAILABLE COPY

C:\Program Files\Sinex\Queries\10521447c.str



chain nodes :
23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 43 44 45 46 47 49 50 52 53
ring nodes :
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22
chain bonds :
1-24 1-52 2-45 3-38 3-39 4-40 4-41 5-23 8-25 10-30 10-31 11-32 11-33 14-26 16-27 17-36 17-37 18-34
18-35 19-46 19-47 20-28 20-29 21-43 22-44 44-50 45-49 52-53
ring bonds :
1-2 1-6 2-3 3-4 4-5 5-6 5-7 6-10 7-8 7-11 8-9 8-14 9-10 11-12 12-13 13-14 13-15 14-18 15-16 15-19
16-17 18-22 17-18 19-20 20-21 21-22
exact/norm bonds :
1-2 1-6 2-3 2-45 3-4 4-5 5-6 5-7 6-10 7-8 7-11 8-9 8-14 9-10 11-12 12-13 13-14 13-15 14-18 15-16
15-19 16-17 16-22 17-18 19-20 20-21 21-22 21-43 22-44 44-50 45-49
exact bonds :
1-24 1-52 3-38 3-39 4-40 4-41 5-23 8-25 10-30 10-31 11-32 11-33 14-26 16-27 17-36 17-37 18-34 18-35
19-46 19-47 20-28 20-29 52-53

G1:H.OH

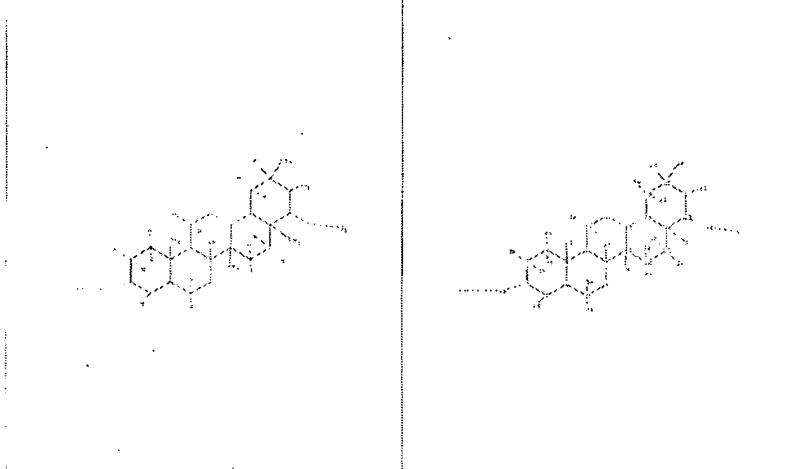
G2:H.Hy

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom 10:Atom 11:Atom 12:Atom 13:Atom 14:Atom
15:Atom 16:Atom 17:Atom 18:Atom 19:Atom 20:Atom 21:Atom 22:Atom 23:CLASS24:CLASS25:CLASS26:CLASS
27:CLASS28:CLASS29:CLASS30:CLASS31:CLASS32:CLASS33:CLASS34:CLASS35:CLASS36:CLASS37:CLASS38:CLASS
39:CLASS40:CLASS41:CLASS43:CLASS44:CLASS45:CLASS46:CLASS47:CLASS49:CLASS50:CLASS52:CLASS53:CLASS

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C:\Program Files\Strnexp\Queries\10521447.str



chain nodes :
23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 43 44 45 46 47 49 50
ring nodes :
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22
chain bonds :
1-24 2-45 3-38 3-39 4-40 4-41 5-23 8-25 10-30 10-31 11-32 11-33 14-26 16-27 17-36 17-37 18-34 18-35
19-46 19-47 20-28 20-29 21-43 22-44 44-50 45-49
ring bonds :
1-2 1-6 2-3 3-4 4-5 5-6 5-7 6-10 7-8 7-11 8-9 8-14 9-10 11-12 12-13 13-14 13-15 14-18 15-16 15-19
16-17 16-22 17-18 10-20 20-21 21-22
exact/norm bonds :
1-2 1-6 2-3 2-45 3-4 3-38 3-39 4-5 4-40 4-41 5-6 5-7 6-10 7-8 7-11 8-9 8-14 9-10 10-30 10-31 11-12
11-32 11-33 12-13 13-14 13-15 14-18 15-16 15-19 16-17 16-22 17-18 17-36 17-37 18-34 18-35 19-20 19-46
19-47 20-21 21-22 21-43 22-44 44-50 45-49
exact bonds :
1-24 5-23 8-25 14-26 16-27 20-28 20-29

G1:H.OH

G2:H.Hy

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom 10:Atom 11:Atom 12:Atom 13:Atom 14:Atom
15:Atom 16:Atom 17:Atom 18:Atom 19:Atom 20:Atom 21:Atom 22:Atom 23:CLASS24;CLASS25;CLASS26;CLASS
27:CLASS28;CLASS29;CLASS30;CLASS31;CLASS32;CLASS33;CLASS34;CLASS35;CLASS36;CLASS37;CLASS38;CLASS
39;CLASS40;CLASS41;CLASS43;CLASS44;CLASS45;CLASS46;CLASS47;CLASS49;CLASS50;CLASS

BEST AVAILABLE COPY

182401, 17

L1 FILE "CARBON" AMENDED AT 09:02:47 ON 02 APR 2007
L2 STRUCTURE UPLOADED
L3 14.3 KB SAM
L4 STRUCTURE UNLINKED
L5 * 0 KB SAM
L6 17.0 KB SAM
L7 PROTECTION UNLINKED
L8 1.0 KB FILE
L9
L10 FILE "CARBON" ENTERED AT 09:08:10 ON 02 APR 2007
L11 * 0 KB
L12 18.0 KB
L13
L14 FILE "RECOVERY" ENTERED AT 09:11:54 ON 02 APR 2007
L15 STRUCTURE UPLOADED
L16 * 0 KB SAM
L17 18.0 KB FILE

RECEIVED